

## SUPPLEMENTAL MATERIAL

### **Identification of Tazarotenic Acid as the First Xenobiotic Substrate of Human Retinoic Acid Hydroxylase CYP26A1 and CYP26B1**

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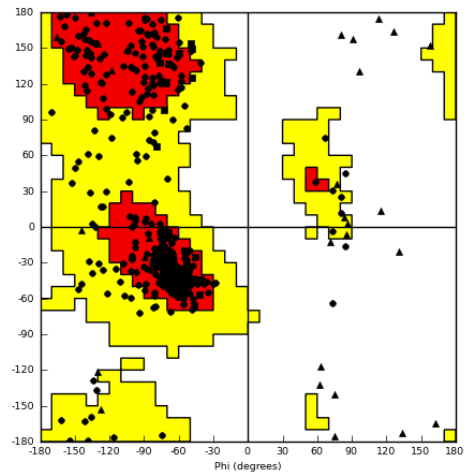
(P.D.)

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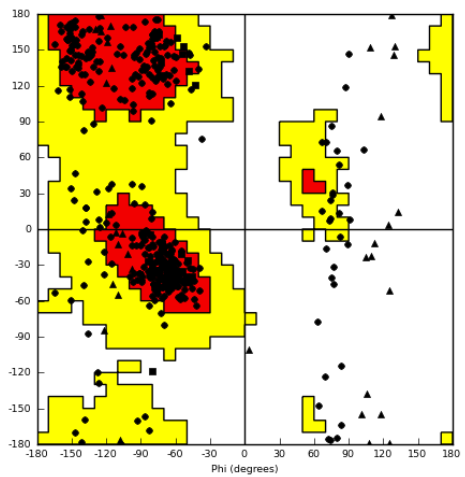
(D.D.)

**Supplemental Figure 1. CYP26A1 and CYP26B1 Ramachandran Plots.** Statistical analysis Ramachandran plots is noted in Table 1.

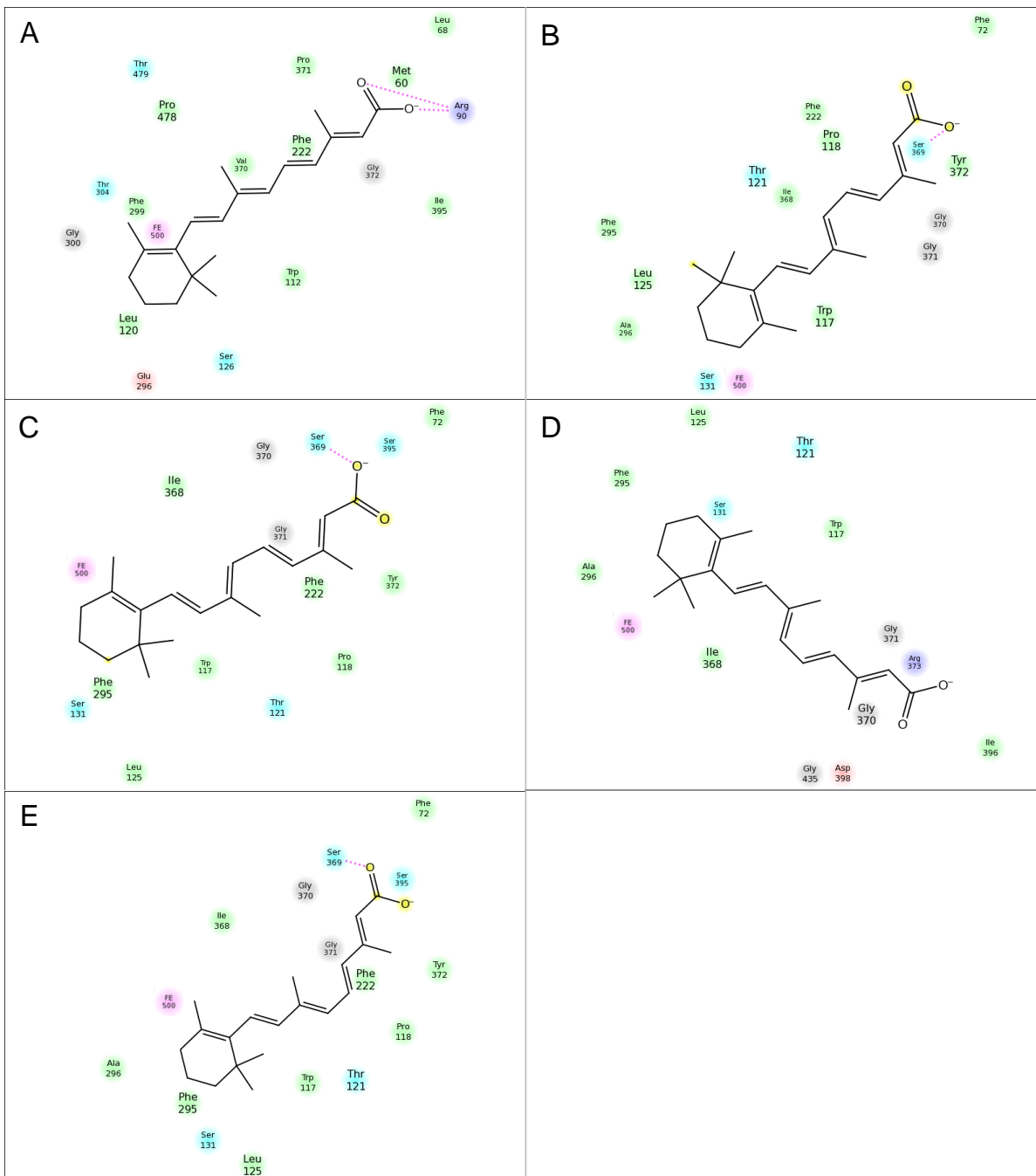
**26A1 Ramachandran Plot**



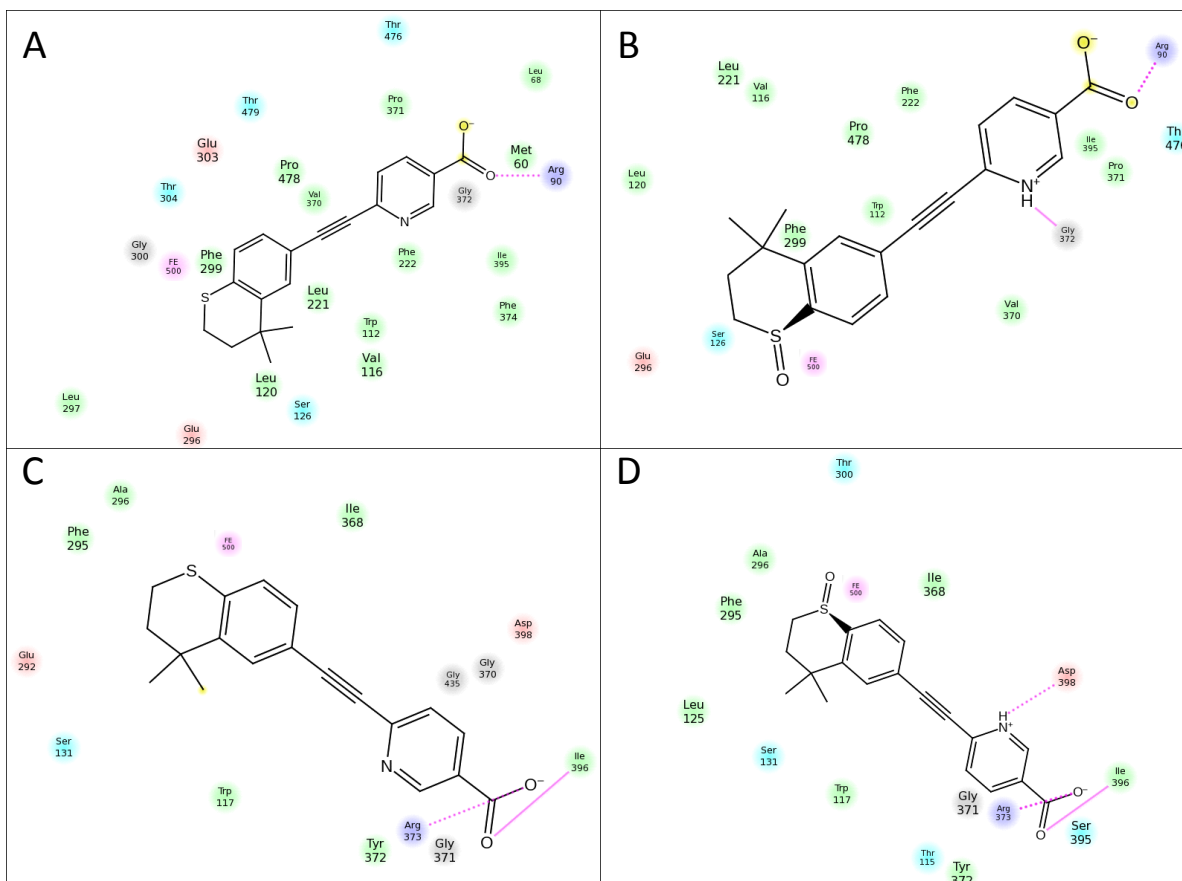
**26B1 Ramachandran Plot**



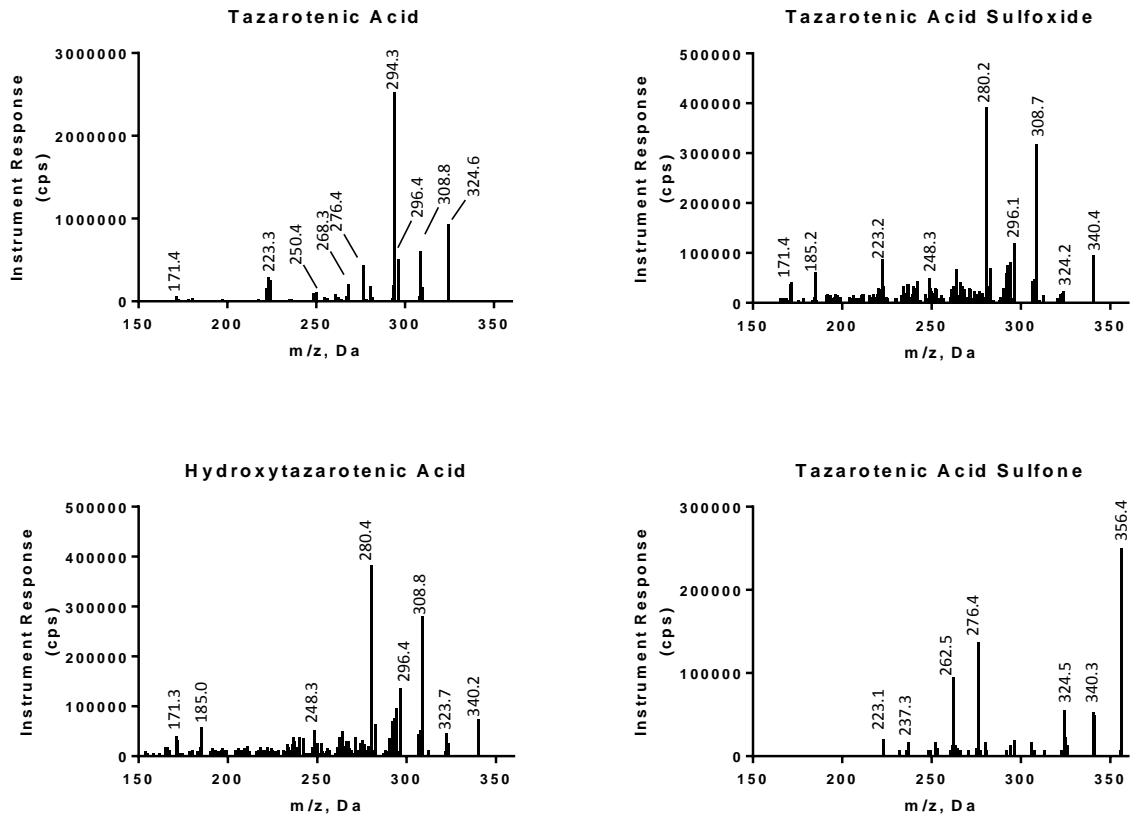
**Supplemental Figure 2. Ligand interaction diagram for *at*-RA docked in the active site of CYP26A1 and CYP26B1.** Analysis of the active site ligand interactions for *at*-RA docked in the active site of CYP26A1 (A) and CYP26B1 (4-(*R*)-OH, B; 4-(*S*)-OH, C; 16-OH, D; 18-OH, E) shows the hydrophobic (green), hydrophilic (blue), electrostatic (red/purple) and metal (pink) interactions located within 3.0 Å of *at*-RA. Hydrogen bonding interactions are depicted by dashed lines.



**Supplemental Figure 3. Figure 6. Ligand interaction diagram for tazarotenic acid or tazarotenic acid sulfoxide docked in the active site of CYP26A1 and CYP26B1.** Analysis of the active site ligand interactions for tazarotenic acid docked in the active site of CYP26A1 (A) and CYP26B1 (C) shows the hydrophobic (green), hydrophilic (blue), electrostatic (red/purple) and metal (pink) interactions located within 3.0 Å of tazarotenic acid. Ligand interactions for tazarotenic acid sulfoxide in the active site of CYP26A1 (B) and CYP26B1 (D) are also shown. Hydrogen bonding is depicted by dashed or solid lines.



**Supplemental Figure 4. MS-MS Spectrum For Tazarotenic Acid (m/z 324.2), Tazarotenic Acid Sulfoxide (m/z 340.3), Hydroxy-Tazarotenic Acid (m/z 340.3) and Tazarotenic Acid Sulfone (m/z 356.3)**



**Supplemental Figure 5. Corresponding Fragmentation Pattern For Tazarotenic Acid (m/z 324.2), Tazarotenic Acid Sulfoxide (m/z 340.3), Hydroxy-Tazarotenic Acid (m/z 340.3) and Tazarotenic Acid Sulfone (m/z 356.3)**

